



"Express Mail" mailing label number EL 856 156 986 US I hereby certify that this document and referenced attachments are being deposited with the United States Postal Service "Express Mail Post Office to Addressee" service under 37 CFR § 1.10 on the date indicated and is addressed to: Commissioner for Patents, Box Patent Application, Washington, D.C. 20231 April 18, 2001 .

By: Printed: Nancy Ramos

### IN THE UNITED STATES PATENT AND TRADEMARK OFFICE

In re Application of: Kaser et al.

Title:

POLYCYCLIC AROMATIC HYDROCARBON INDUCED MOLECULES

Serial No.:

To Be Assigned

Filing Date:

Herewith

Examiner:

To Be Assigned

Group Art Unit:

To Be Assigned

Official Draftsman

Commissioner for Patents Washington, D.C. 20231

### SUBMISSION OF FORMAL DRAWINGS

Sir:

Transmitted herewith are Figure(s) 1A, 1B, 1C, 1D, 1E, 1F, 2A, 2B, 2C, 3, 4, as Eleven (11) sheets of formal drawings for this application. Each sheet of drawing indicates the identifying indicia suggested in 37 CFR Section 1.84(c) on the reverse side of the drawings.

Applicants believe that no fee is due with this paper. However, if the Commissioner determines that a fee is necessary, the Commissioner is hereby authorized to charge any additional fees associated with this communication or credit any overpayment to Deposit Account No. 09-0108. A duplicate copy of this communication is enclosed.

If there are any questions regarding the above, the Examiner is invited to call the undersigned at 650-855-0555.

Respectfully submitted,

INCYTE GENOMICS, INC.

Date: 18h April 2001

Matthew R. Kaser, D.Phil.

Reg. No. 44,817

Direct Dial Telephone: (650) 845-4596

3160 Porter Drive Palo Alto, California 94304 Phone: (650) 855-0555

Fax: (650) 845-4166

TCA 110 GCT 164 119 TTG AAG CAA TTA AAC GTT CAA ATG AAT ATG AGT AAT GTA ATG GGA M N M S N V M G 5' CTTCA GTT CCT CGA AGA GAT TCA CTT TCA AAA ACA TCA ACT CCT TAT AAC AAA GTG TTT 65 74 83 92 101 GCA GCA AGC CAA GGG ACC CCA TGG GAA ACA CTT GTC AAA AAC

AAT

218 AGT S 272 TCA GTA GGA S V G CAA TTA Q TCA 263 GGA AGA ' GGT TTG AAG AGC CAG GGC CGT CTG G L K S Q G R L CTG L 200 254 CGA GAG ATC AGC ATG TCT GTT GGT R E I S M S V G 191 ACA ACT TGG ACA ACT AGT T T W T T S 182 236 GATQ CGT AAT z

GAT D 326 GTTGTC AAT GCT TTG GAG ATG V N A L E M 317 308 GTA GTT GGA GGG ACC ATA GAT V V G G T I D 299 290 GGA G GGA 281 AAA K TCT S

380 ACT T CCA AAT CAG CAA CCC AGT CAC AAA ATT CAG ATT P N Q Q P S H K I Q I 371 362 353 GAA CAT (E H 344 ATT TCT ( I S ) CAT

9 9 ATG CTC. TCT ACT GAA GCT CGT GTT GAT TAC ATG GGC TCA AGT ATC S T E A R V D Y M G S S I 416 407 398 ATG GGT N

DSESOUL THISSE AAT GCT GAT TTC

CCA GAT CTG
P D L GAT650 AGC ATG 704 GAA CAA TTA CTT GAT GCA GCA CAT CAT E Q L L D A A H H 758 ഗ  $\operatorname{TTT}$ GGA G TGG AAA GTA AAC TTG TTT TTC ACA CAA TTT GAT ACC F F T Q Q F D T CTT CCG CCA AAG ACA L P P K T TGC CAC ATA TCC TTA C H I S L CAT GGA GGA TCA ATG AGC G G S M S GIC CAA GTA ATG ATA TCA AGG TCA ACC ACA Q V M I S R S T T 533 > 587 641 695 749 857 ATT TTC (I F M CAG GAT GAA Q D E GAT TCA AGC ATA ACT GAT AAA AGT GAG D S S I T D K S E 686 GTT CCT TAC ( V P Y I 578 632 CCT GGA GTA TTG AAG GTG GTA TCA GGA P G V L K V V S G 740 794 848 GAA GAT GGA ATG CAA TTT 515 GAA E CCA P 569 CAA O 623 677 731 785 CTT AAG ( L K I GGA ATG AAG CTC CAG G M K L Q CGA GCT CTG TCT ACC TGG GGA R A L S T W G CTA GAA AAA AGT TCA L E K S S TTC 506 Д 560 614 899 722 212 830 ATTTTA CCA ( L P Ø Z Д AGT  $ext{TTG}$ ATA AAA ATA ACT AGC AAC (T S N ] CGA CAC TGG (R H M 605 629 AAG ATT I ATC

920 TCT	974 CAG O	)28 )TA	.82 .AA	36 CC
AAA K	GCT (	10 ACA C	_ 1082 TTG GAA L E	11 CA C
TCA AAA S K	GAA ( E	'AA	GCT T	1127 CAT GAA AAT CCA CCC H E N P P
911 CGT R	947 CCA AAT ATT GCT TTT TGG ACT ( P N I A F W T I	019 GTA (	073 3GT G	.27 JAA A
TTT F	TGG W	1 ATT ( I	1073 TGT GGT C	11 'AT G
AAT N	TTT F	TAT	CCA 7	, dc
902 CCA	956 GCT A	1010 ACA T	1055 AAT ACT ATG GTT ACC AAA N T M V T K	1109 AAG ATA ACA AGG CGT CGC ( K I T R R R I
G G	ATT I	TCT S	1 ACC T	1 AGG ( R )
893 TTT CAT ( F H (	AAT N	ZAT I	GTT V	ACA
893 TTT F	947 CCA P	1001 3 AGT GAT C	.055 ATG M	109 ATA I
TGT C	GAA	AGT	ACT T	1 AAG K
g GCA	GAA	TCC	AAT N	JCC.
884 A CTG L	938 TTA L	992 GGC G	1046 CTG GGT CAT ; L G H 1	1100 GCA ACA ATA A A T I T
ATG ACA ( M T	' CAT H	GAA GAT E D	GGT	1 ACA T
F ATC	TTT (	GAA E	CTG L	GCA A
875 CAT H	929 CTT L	983 TGG W	1037 CAC H	1091 ' ATG M
GGA AAT G N	TGG GCC W A	983 AAA ATC TGG K I W	TTT F	CCT P
GG2	TGG ¥	AAA K	1037 GAT TTT CAC ( D F H I	1091 AGT CCT ATG G S P M A

ACT GTG AAG AAT TCT AGT TTG TTG AGT GGA TTC AGA GGA GGT TCT AGC TAC AAC
T V K N S S L L S G F R G G S S Y N
FIGURE 1C

CAT GGA GTA GCA AGT GTG AAA GAA TGG TTC AAT TAT GTT ACA GCT ACA AGG AAT
H G V A S V K E W F N Y V T A T R N

1172

1163

1154

1145

GAA GAG CTA AAT CTG CTT CGT AAT GTT GAT GCT AAC AAC ACT GAG AAT AGC ACT E L N L L R N V D A N N T E N S T

1253

CAT GAA ACA GAG ACT ATC TTT GCA TTA CCA AGG ATG CAG CTT GAC TTT AAA TCC H E T E T I F A L P R M Q L D F K S 

ATT CAT GTT CAA GAA CCA CAG GAG CCT TCA TTA CAG GAT GCC AGC CTG AAG CCA I H V Q E P Q E P S L Q D A S L K P 

SUBCLASS

AAA GTA GAA TGT AGT GTG ACA GAG TTC ACT GAC CAC ATT TGT GTG ACT ATG K V E C S V V T E F T D H I C V T M 

GAT GCT GAG CTC ATC ATG TTT CTT CAT GAT TTA GTA TCA GCT TAT CTT AAA GAA D A E L I M F L H D L V S A Y L K E 

AAA GAA AAA GCC ATC TTT CCA CCT CGG ATT TTA TCT ACT CGA CCA GGA CAA AAA K E K A I F P P R I L S T R P G Q K 

AGT CCA ATT ATA CAT GAC GAC AAT TCC TCT GAT AAA GAT AGA GAA GAT AGC S P I I H D D N S S D K D R E D S 

ATC ACT TAT ACT ACT GTG GAC TGG AGA GAT TTT ATG TGC AAT ACA TGG CAC CTA I T Y T T V D W R D F M C N T W H L 

### FIGURE 1D

GAA CCT ACT CTT AGA TTA ATT TCT TGG ACT GGA AGA AAG ATT GAT CCA GTA GGT E P T L R L I S W T G R K I D P V G

GTT GAT TAT ATT CTT CAA AAA TTG GGC TTT CAT CAT GCT AGG ACT ACT ATT CCT V D Y I L Q K L G F H H A R T T I P 

AAA TGG CTT CAA AGA GGA GTC ATG GAT CCA CTG GAC AAG GTT CTG TCA GTT CTT K W L Q R G V M D P L D K V L S V L 

ATC AAA AAG CTC GGT ACT GCA CTA CAG GAT GAA AAG GAA AAG AAA GGC AAA GAC I K K L G T A L Q D E K E K K G K D 1865 1874

AAA GAA GAA CAC TAA AAA AGT AAT TTG ATC TGT GAA CAA ATT ATG ATT GTG TCT 

GTT TTA TTA CAC TGG AGT GTT TTT TTA GTA TAA TAA TTT GAA ATA TAA CTT TAA 

AAT AAT TCT AAA TTT GTG GCT ATA ATT AAA AGT TTG TAA GTT AAC CTG TTC TAG 

## FIGURE 1E

# ISSISSIL INTESSI

TTC CAT CAT TCT GTG TAC AGT GAA GTA TTG CAT GAT AAT GTA AAT TTT GTG AAA 

AAC TAG ATT AAA ATA TAT AAC TGC TTG TTA TGG TTT ATA ATT ATA TAT TGT GCA 

SUBCLASS

ATA CAA TIC CIG CAI CIT IAA AAI GIC IGC AGA AIA ACI GIG AAI III III GII 

ATT GGA TTG GCC GTA ACT TTT AGA AAA AAA TCT TGT TGA TGA TAA TGT GAT TTT 

GGG GAG GTC AIT AAT TGC TTT TTT TTT AAA TGT AGA CTT ATA TAA ATA CCT 

GTT TGT ATA TAG CTT GAG TAA TTG TGA TAT GAT TGT ATA CCA CTA AAA TAT TGT 

AAA AAA A 3'

## FIGURE 1F

SUBCLASS 47

56 GTG V CAT H CCC ( GTG V GTG GAC (V D ) 5' GCTTG AAG GCG GGA AAA TAC GGA ATG GAG CGA CAG GGA

110 CGG R CAT CCG (H P 1 GAG (E 101 AAG K AGC 2 CCG CAG CCC TTG GGG GAG
P Q P L G E CAG GAA ( Q E ] GAC (D AAA K TGC C AAA K

164 TGC C 137 146 155 GCT GGT GGA GGG CCA GCT AGT GCC AGT A G G G P A S A S 128 3 TGC GAG GAG GAA G C E E E E GAG AAC TE N GAA E

218 TCA S 209 TTC TCC TCT CTG GAC F S S L D 200 TAC ' Y 191 2 TCG GGA CTC 1 S G L 3 GGG AAG G 173 182 CTG ACG GTC CTG GAA L T V L E 173 CAG Q

272 AAG K AAC N ATC I 254 263 F CTG ATC GAA AAC A L I E N I CAG GAG (Q 245 GCC ( 236 ATC CTG CAG AAG AGA I L Q K R GAC 1 227 ATT

CTG AAG L 290 299 308 317
CAT GCA CTC ATG ACC AAC TTC AGG AAC AGC H A L M T N F R N S 290 AAG GAC ( K D I

### FIGURE 2A

AAT N TAT Y ATT CAG Q ATC TAT ( I Y ( 362 GAG AGG 1 E R GAG E 353 GAG AAA TTA ( E K L I ACA GAT CTG 335 GTT TCG ( V S I AAG K

434 GCA A CAG AAA ATG O 416 GAG TTC ACC ( E F T ( CTC CAA ( 407 CAG GAA AAG C Q E K I ATC AAG ATC ; K I AAC N 389 CAC H GAC D

488 GTG GAG ACT V E T AGC S 479 TGC CAC ACA GAG CTC AAA CAA GTC T E L K Q V 470 461 GAG . 452 TTG CAT H 443 AGC S ATC I

542 GGG G AGA TGG R W 533 CTA AGA CTC A L R L F 524 GAG CAG AGC ( E Q S I 515 CCT P 7 4 GAC CTG TGT CTC CAG C D L C L Q F AAA K 497 GTG TAC V

596 CCA P CCC P CAG Q 587 GGC AAC TCA C G N S C 578 CCA CGT CCC ( 569 CCA P AAG TCC K 560 TCT AGG GGA A S R G K 551 CAC H GAC D

650 GAA E TCA S GTT TCT TCT GTG GCT GAA ACT ACT TCT CAG GCC ACT GCT V S S V A E T T S Q A T A 605 TTC F

### FIGURE 2B

GTA CAG ACC AAC AGA GAT GGT GAA TGC TGA CAG CTG CCG GGA GAC TCA CGC CTT V Q T N R D G E C 989

13 ISUBCLASS

AGT GAC AGT CTC CAG GAG AAG ACT GTG AGG CCA CCA TTT GGG CCA CAC TGA GAA 740 731

ATT GIT TIT CAT GGT TCT ATA ATG CAT CTT GGC AGA AAA AAA CAA AAA CCC AAA 794 785

CCC GCT CCT TGT GCT GAA CTC CCA AAA TGT AGC AAG TCC AGC CCT CTC CAT AGG 848 839 830

AGG CTT CGT GCT CCC CAC CCT TGG CAA GTT CTC CCC ACC CCC AGC CCC ACA GTT 902 893 884

929 TAT TAA ATG TTT GAT TTT CAA AAA 3

### FIGURE 2C

IG.

TCC S AGT S CTG ACC I 5' CTC AAG ACC CAG TGG GAC AGC CAG ACA GAC GGC ACG ATG GCA CTG AGC GCC GCT TGC CTC CTC CTC CTC GCC AGC A A C L L L L L L L L S S ATC TGG (I M

TTC CCA CAA CAG GGA CAA CTT GCA GAG CTG CAA CCC CAG GAC F P Q Q T G Q L A E L Q P Q D 153 144 126 GTT ' TCT (S 999 1990

CCC ATG TTC CAG AGG CGA AGG AGG CGA P M F Q R R R R R 216 207 198 AGA GCT GGA GCC AGC TGG ATG R A G A R A S W M 189 180

CGA TCA AAG R S K GGC TGC TGT CAT G C C H 252 GAC ACC CAC TTC CCC ATC TGC ATT TTC TGC TGC D T H F P I C I F C C

324 ပ္ပ TGC TGC AAG ACG TAG AAC CTA CCT GCC CTG CCC CCG TCC CCT ATG ' 999

TTG GAA TAA AAT GGC 369 333 342 351 360 ATT TAT TCC TGC TGC CCC AGA ACA TAG GTC CTT TTC

387 TTC TTT TGT TTT CCA AAA AAA AA 3' FIGURE 3

DRAFTS:

MALSSQIWAACLLLLLLASLTSGSVFPQQ 1642580 MALSTRIQAACLLLLLLASLSSGAYLRQQ 700139271

TGQLAELQPQDRAGARASWMPMFQ-RRRRR 1642580 TRQTTALQPWHGAESKTDDSALLMLKRRKR 700139271 전 8 3

60 DTHFPICIFCCGCCHRSKCGMCCKT 60 DTNFPICLFCCKCCKNSSCGLCCIT

1642580 700139271

FIGURE 4